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Molecular mechanisms of desiccation tolerance in the resurrection glacial relic *Haberlea rhodopensis*

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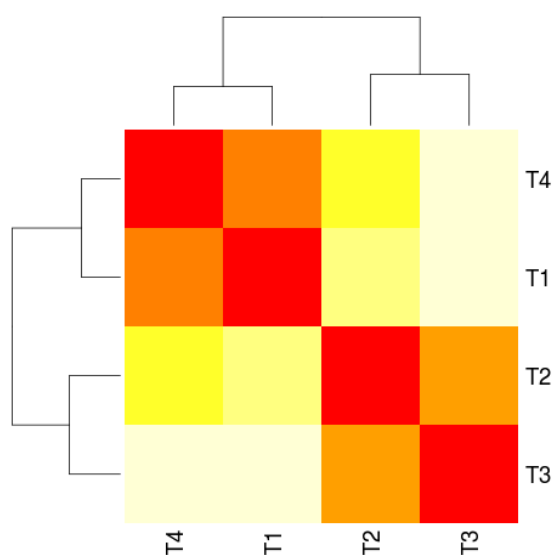
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Supplemental Figure 1. Cluster analysis of the four growth conditions based on the next generation sequence (NGS) transcriptome data. T1, unstressed controls; T2, drought stress; T3, desiccation; T4, rehydration. The intensity of the colour corresponds to similarity in gene expression. Unstressed controls cluster with rehydrated plants (T1 & T4) while drought stressed and desiccated plants cluster together (T2 & T3).